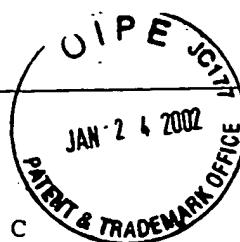


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Fanslow III, William C

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A10

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Thr Ile Tyr Phe Arg Asp Phe Arg Pro Leu Lys Asp Ser Arg Phe Gln
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A10

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A10 cat gaa tct cta aag gga gct gac aga tcc acc tct caa cgc att gaa 942
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A10

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Arg Glu Lys Asp Thr Ala Thr Leu Asn Cys Gln Ser Ser Gly Ser Lys
180 185 190

Pro Ala Ala Arg Leu Thr Trp Arg Lys Gly Asp Gln Glu Leu His Gly
195 200 205

Glu Pro Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val
210 215 220

Ser Ser Ser Val Thr Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Ser
225 230 235 240

Ile Val Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser
245 250 255

Thr Ser Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg
260 265 270

Pro Asp Pro Pro His Pro Arg Glu Gly Gln Lys Leu Leu Leu His Cys
275 280 285

A10
Glu Gly Arg Gly Asn Pro Val Pro Gln Gln Tyr Leu Trp Glu Lys Glu
290 295 300

Gly Ser Val Pro Pro Leu Lys Met Thr Gln Glu Ser Ala Leu Ile Phe
305 310 315 320

Pro Phe Leu Asn Lys Ser Asp Ser Gly Thr Tyr Gly Cys Thr Ala Thr
325 330 335

Ser Asn Met Gly Ser Tyr Lys Ala Tyr Tyr Thr Leu Asn Val Asn Asp
340 345 350

Pro Ser Pro Val Pro Ser Ser Ser Thr Tyr His Ala Ile Ile Gly
355 360 365

Gly Ile Val Ala Phe Ile Val Phe Leu Leu Leu Ile Met Leu Ile Phe
370 375 380

Leu Gly His Tyr Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu
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gccagcgccc agccagggag ccggccggga agcgcg atg ggg gcc cca gcc 174
Met Gly Ala Pro Ala Ala
1 5
tcg ctc ctg ctc ctg ctg ttc gcc tgc tgc tgg gcg ccc ggc 222
Ser Leu Leu Leu Leu Leu Leu Phe Ala Cys Cys Trp Ala Pro Gly
10 15 20
ggg gcc aac ctc tcc cag gac gac cag ccc tgg aca tct gat gaa 270
Gly Ala Asn Leu Ser Gln Asp Asp Ser Gln Pro Trp Thr Ser Asp Glu
25 30 35
aca gtg gtg gct ggt ggc acc gtg gtg ctc aag tgc caa gtg aaa gat 318
Thr Val Val Ala Gly Gly Thr Val Val Leu Lys Cys Gln Val Lys Asp
40 45 50
cac gag gac tca tcc ctg caa tgg tct aac cct gct cag cag act ctc 366
His Glu Asp Ser Ser Leu Gln Trp Ser Asn Pro Ala Gln Gln Thr Leu
55 60 65 70
tac ttt ggg gag aag aga gcc ctt cga gat aat cga att cag ctg gtt 414
Tyr Phe Gly Glu Lys Arg Ala Leu Arg Asp Asn Arg Ile Gln Leu Val
75 80 85
acc tct acg ccc cac gag ctc agc atc agc atc agc aat gtg gcc ctg 462
Thr Ser Thr Pro His Glu Leu Ser Ile Ser Ile Ser Asn Val Ala Leu
90 95 100
gca gac gag ggc gag tac acc tgc tca atc ttc act atg cct gtg cga 510
Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile Phe Thr Met Pro Val Arg
105 110 115
act gcc aag tcc ctc gtc act gtg cta gga att cca cag aag ccc atc 558
Thr Ala Lys Ser Leu Val Thr Val Leu Gly Ile Pro Gln Lys Pro Ile
120 125 130
atc act ggt tat aaa tct tca tta cgg gaa aaa gac aca gcc acc cta 606
Ile Thr Gly Tyr Lys Ser Ser Leu Arg Glu Lys Asp Thr Ala Thr Leu
135 140 145 150
aac tgt cag tct tct ggg agc aag cct gca gcc cgg ctc acc tgg aga 654
Asn Cys Gln Ser Ser Gly Ser Lys Pro Ala Ala Arg Leu Thr Trp Arg
155 160 165
aag ggt gac caa gaa ctc cac gga gaa cca acc cgc ata cag gaa gat 702
Lys Gly Asp Gln Glu Leu His Gly Glu Pro Thr Arg Ile Gln Glu Asp
170 175 180
ccc aat ggt aaa acc ttc act gtc agc agc tcg gtg aca ttc cag gtt 750
Pro Asn Gly Lys Thr Phe Thr Val Ser Ser Ser Val Thr Phe Gln Val
185 190 195

A10

acc cg_g gag gat gat ggg g_c agc atc gt_g t_gc t_ct gt_g aac cat gaa 798
 Thr Arg Glu Asp Asp Gly Ala Ser Ile Val Cys Ser Val Asn His Glu
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t_ct c_ta a_ag g_ga g_ct g_ac a_ga t_cc a_cc t_ct c_gc a_tt g_aa g_tt t_ta 846
 Ser Leu Lys Gly Ala Asp Arg Ser Thr Ser Gln Arg Ile Glu Val Leu
 215 220 225 230

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 Tyr Thr Pro Thr Ala Met Ile Arg Pro Asp Pro Pro His Pro Arg Glu
 235 240 245

g_gc c_ag a_ag c_tg t_tg c_ta c_ac t_tg g_ag g_gt c_gc g_gc a_at c_ca g_tc c_{cc}c 942
 Gly Gln Lys Leu Leu Leu His Cys Glu Gly Arg Gly Asn Pro Val Pro
 250 255 260

c_ag c_ag t_ac c_ta t_gg g_ag a_ag g_gc a_gt g_tg c_ca c_{cc}c c_tg a_ag a_tg 990
 Gln Gln Tyr Leu Trp Glu Lys Glu Gly Ser Val Pro Pro Leu Lys Met
 265 270 275

acc c_ag g_ag a_gt g_cc c_tg a_tc t_tc c_ct c_tc a_ac a_ag a_gt g_ac a_gt 1038
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 280 285 290

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 Tyr Tyr Thr Leu Asn Val Asn Asp Pro Ser Pro Val Pro Ser Ser Ser
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 Ser Thr Tyr His Ala Ile Ile Gly Gly Ile Val Ala Phe Ile Val Phe
 330 335 340

ct_g ct_g c_tc a_tc a_tg c_tc a_tc t_tc c_tt g_gc c_ac t_ac t_tg a_tc c_{gg}g c_ac 1230
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 345 350 355

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 Lys Gly Thr Tyr Leu Thr His Glu Ala Lys Gly Ser Asp Asp Ala Pro
 360 365 370

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 375 380 385 390

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35 40 45

Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp Ser Asn
50 55 60

Pro Ala Gln Gln Thr Leu Tyr Phe Gly Glu Lys Arg Ala Leu Arg Asp
65 70 75 80

Asn Arg Ile Gln Leu Val Thr Ser Thr Pro His Glu Leu Ser Ile Ser
85 90 95

Ile Ser Asn Val Ala Leu Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile
100 105 110

Phe Thr Met Pro Val Arg Thr Ala Lys Ser Leu Val Thr Val Leu Gly
115 120 125

Ile Pro Gln Lys Pro Ile Ile Thr Gly Tyr Lys Ser Ser Leu Arg Glu
130 135 140

Lys Asp Thr Ala Thr Leu Asn Cys Gln Ser Ser Gly Ser Lys Pro Ala
145 150 155 160

Ala Arg Leu Thr Trp Arg Lys Gly Asp Gln Glu Leu His Gly Glu Pro
165 170 175

Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val Ser Ser
180 185 190

Ser Val Thr Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Ser Ile Val
195 200 205

Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser Thr Ser
210 215 220

Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg Pro Asp
225 230 235 240

Pro Pro His Pro Arg Glu Gly Gln Lys Leu Leu Leu His Cys Glu Gly
245 250 255

Arg Gly Asn Pro Val Pro Gln Gln Tyr Leu Trp Glu Lys Glu Gly Ser

A10

260

265

270

Val Pro Pro Leu Lys Met Thr Gln Glu Ser Ala Leu Ile Phe Pro Phe
275 280 285

Leu Asn Lys Ser Asp Ser Gly Thr Tyr Gly Cys Thr Ala Thr Ser Asn
290 295 300

Met Gly Ser Tyr Lys Ala Tyr Thr Leu Asn Val Asn Asp Pro Ser
305 310 315 320

Pro Val Pro Ser Ser Ser Thr Tyr His Ala Ile Ile Gly Gly Ile
325 330 335

A10
Val Ala Phe Ile Val Phe Leu Leu Ile Met Leu Ile Phe Leu Gly
340 345 350

His Tyr Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu Ala Lys
355 360 365

Gly Ser Asp Asp Ala Pro Asp Ala Asp Thr Ala Ile Ile Asn Ala Glu
370 375 380

Gly Gly Gln Ser Gly Gly Asp Asp Lys Lys Glu Tyr Phe Ile
385 390 395
